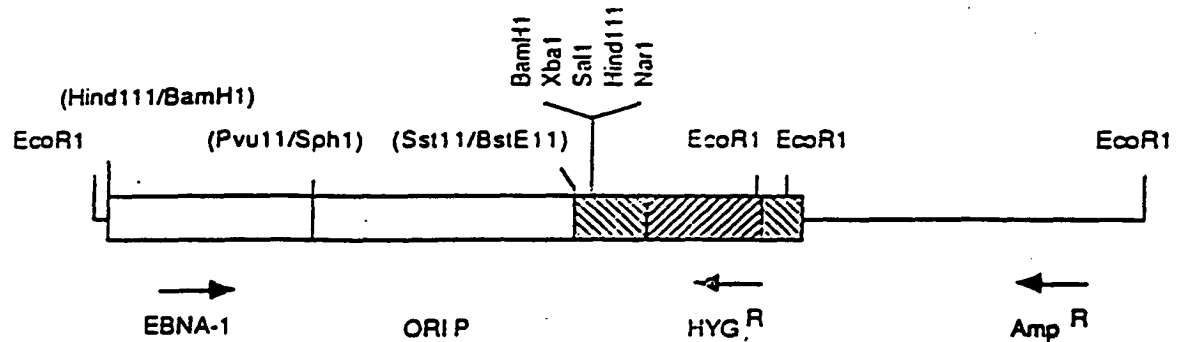


# Figure 1 EBV-Based Self-Replicating Expression Vector

**p220.2** is a 8952 bp plasmid which encodes for EBNA-1, OriP and Hygromycin resistance. It replicates as plasmid in 143 and HeLa cells. EBNA-1 in this construct is driven off an unknown promoter located in the pBR322 sequences. DNA inserted upstream of EBNA-1 appears to eliminate expression of EBNA-1.



bp

1-35

— pBR322

36-2646



EBV EBNA-1 107567-110176 (Baer et. al., Nature 310:1954) Bam H1-Pvu11 fragment. Bam H1 site was blunt-end ligated to the Hind111 site.

2647-4826



EBV OriP 7333-9516 Sph1- Sst11 sites blunt-end ligated to the BstE11 site. (Sugden et.al., MCB 5: 410, 1985)

4827-5460

6488-6747



HSV TK regulatory region (McKnight, S.L., Nucleic Acids Res. 8, 5949, 1980.) Pvu11 fragment ligated into the poisonless pBR322 at Nae I site. These sites lost in cloning.

5461-6487



HPH gene (Gritz and Davies, Gene 25:179, 1983) Ban H1 fragment blunt end ligated into the Sma1 and Bgl1<sup>+</sup> sites in HSV TK sequences.

6748-8952

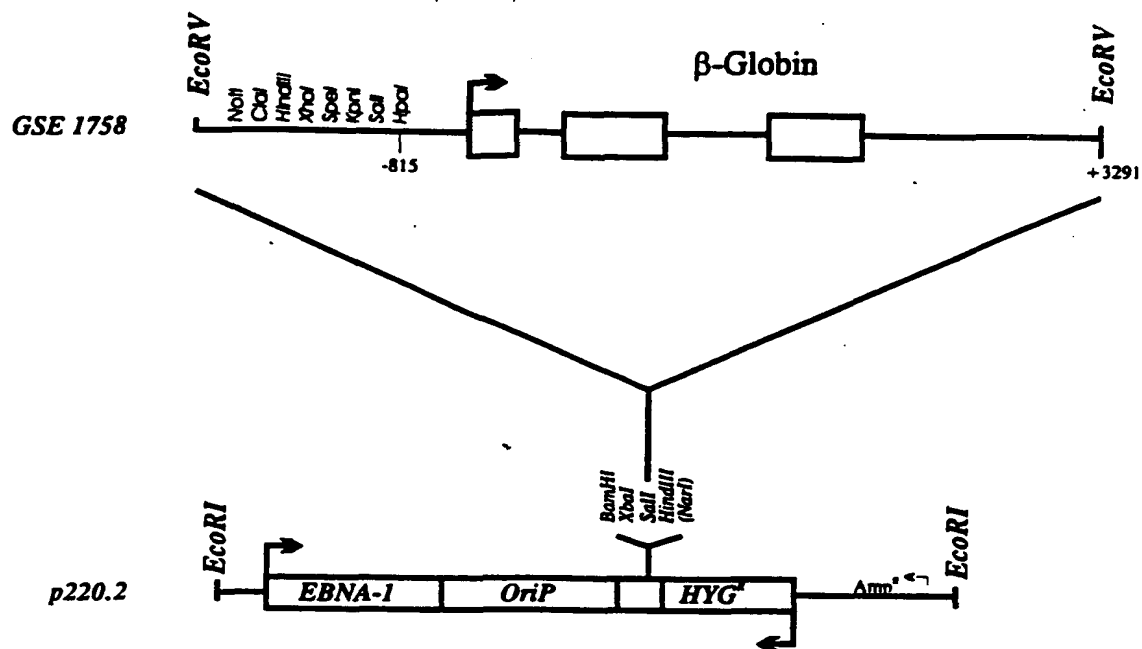


pBR322 poisonless vector (deletion of 1.1 kb in pBR322) confers ampicillin resistance. (Lusky & Botchan, Nature 293:79,1981)

The polylinker from pUC 12 (Sma1-Hae111 fragment) is inserted into a Nar1 site within the HSV TK sequences. The Pst1 site in the polylinker is not unique.

(/) denotes "blunt-end ligations", these sites were not regenerated in cloning.

0947054-15024260



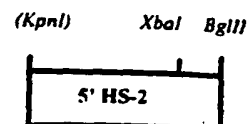
**Figure 2: Reporter gene construct.**

The  $\beta$ -globin gene extending from a 5' *HpaI* site at -815bp to an *EcoRV* site 1685bp passed the poly(A)-addition site in the plasmid GSE1758 (Talbot et al., 1990) was removed as a 4.1kb *EcoRV* fragment and inserted into a blunted *SalI* site in the polylinker of p220.2 (Figure 1). Note: this cloning step brings a number of extra restriction enzyme sites (including a unique *SalI* site) 5' of the  $\beta$ -globin gene.

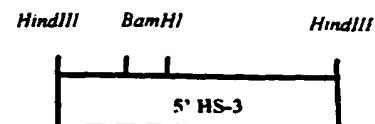
**Reference:**

Talbot, D., Philipsen, S., Fraser, P. and Grosveld, F. (1990) EMBO J. 9: 2169-2178.

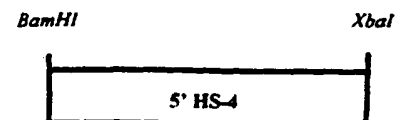
5' HS-2 - 1.5kb KpnI-BglII  
blunted fragment



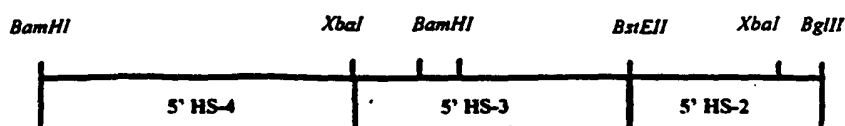
5' HS-3 - 1.9kb HindIII  
fragment



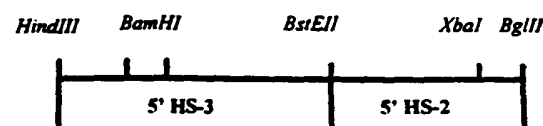
5' HS-4 - 2.1kb BamHI-XbaI  
fragment



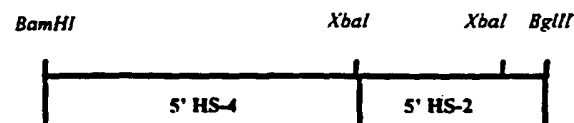
5' HS-4-3-2  
5.5kb construct



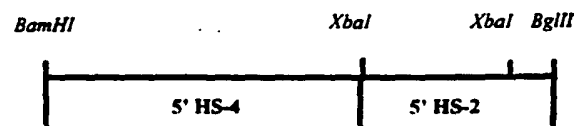
5' HS-3-2  
3.4kb construct



5' HS-4-3  
4kb construct

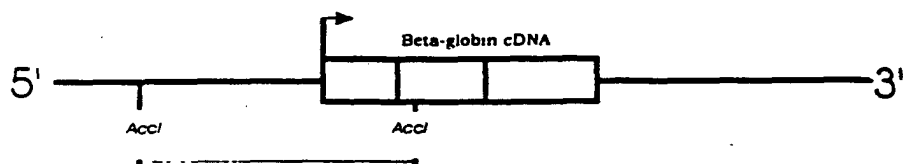
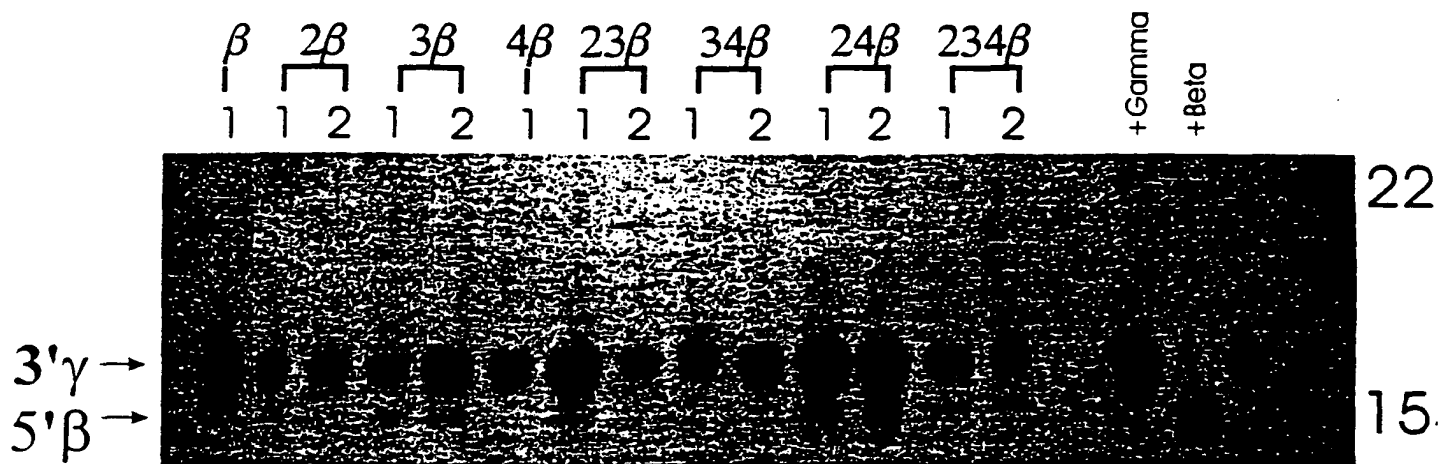


5' HS-4-2  
3.6kb construct

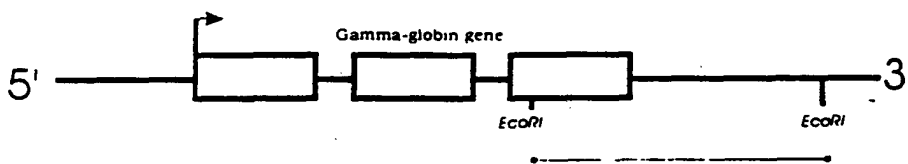


**Figure 3:  $\beta$ -Globin LCR hypersensitive site constructs**

Multiple hypersensitive site constructs retained the site order found in the wild type  $\beta$ -globin locus. *SalI* linkers were added to both the 5' and 3' ends allowing the DNA to be cloned into the unique *SalI* site in the  $\beta$ -globin-p220.2 reporter vector (Figure 2).



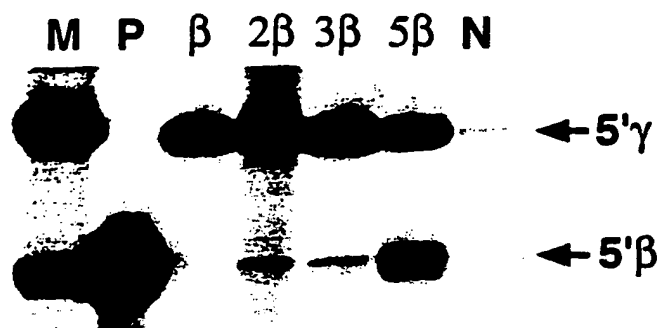
155nt S1-protected fragment



168nt S1-protected fragment

Figure 4: Numbers represent  $\beta$ -globin locus control region DNaseI hypersensitive site combination used.

# A. K562



# B. HeLa

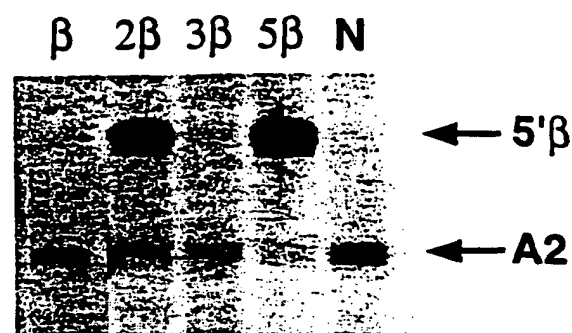


Figure 5.

Fig. 6

Expression Analysis of  $\beta$ LCR/Episome Constructs in K562 Cells

